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TECH CENTER 1600/2900

RAW SEQUENCE LISTING

DATE: 12/10/2001

PATENT APPLICATION: US/09/848,798A

TIME: 11:15:20

Input Set : N:\Crf3\RULE60\09848798A.txt

Output Set: N:\CRF3\12102001\I848798A.raw

5 <110> APPLICANT: Siegel, Donald L.
 9 <120> TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
 11 SORTING METHOD FOR PRODUCTION THEREOF
 15 <130> FILE REFERENCE: 09596-42U2
 19 <140> CURRENT APPLICATION NUMBER: 09/848,798A
 20 <141> CURRENT FILING DATE: 2001-05-04
 24 <150> PRIOR APPLICATION NUMBER: 09/240,274
 25 <151> PRIOR FILING DATE: 1999-05-02
 28 <150> PRIOR APPLICATION NUMBER: 60/028,550
 30 <151> PRIOR FILING DATE: 1996-10-11
 34 <160> NUMBER OF SEQ ID NOS: 224
 38 <170> SOFTWARE: PatentIn Ver. 2.0
 42 <210> SEQ ID NO: 1
 44 <211> LENGTH: 128
 46 <212> TYPE: PRT
 48 <213> ORGANISM: Homo sapiens
 52 <220> FEATURE:
 54 <223> OTHER INFORMATION: anti-Rh(D) chain B01
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 66 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Arg Ser Tyr
 68 20 25 30
 72 Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 74 35 40 45
 78 Ala Ala Thr Ala Tyr Asp Gly Lys Asn Lys Tyr Tyr Ala Asp Ser Val
 80 50 55 60
 84 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Phe
 86 65 70 75 80
 90 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Phe Tyr Cys
 92 85 90 95
 96 Ala Arg Gly Gly Phe Tyr Tyr Asp Ser Ser Gly Tyr Tyr Gly Leu Arg
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 136 1 5 10 15
 140 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe Ser Ser Tyr
 142 20 25 30
 146 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

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148          35          40          45
152 Ser Val Ile Ser Tyr Asp Gly His His Lys Asn Tyr Ala Asp Ser Val
154          50          55          60
158 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
160 65          70          75          80
164 Leu Gln Met Asn Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys
166          85          90          95
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210          20          25          30
214 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
216          35          40          45
220 Ser Val Ile Ser Tyr Asp Gly His His Lys Asn Tyr Ala Asp Ser Val
222          50          55          60
226 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
228 65          70          75          80
232 Leu Gln Met Asn Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys
234          85          90          95
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278          20          25          30
282 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
284          35          40          45
288 Ser Val Ile Ser Tyr Asp Gly His Asn Lys Asn Tyr Ala Asp Ser Val
290          50          55          60

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294 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
296 65 70 75 80
300 Leu Gln Met Asn Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys
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346 20 25 30
350 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
352 35 40 45
356 Ala Val Ile Ser Tyr Asp Gly Thr Asn Lys Tyr Phe Ala Asp Ser Val
358 50 55 60
362 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
364 65 70 75 80
368 Leu Gln Met Thr Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Phe Cys
370 85 90 95
374 Ala Asn Leu Arg Gly Glu Val Thr Arg Arg Ala Ser Val Pro Leu Asp
376 100 105 110
380 Ile Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser
382 115 120
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412 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe Ser Ser Tyr
414 20 25 30
418 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
420 35 40 45
424 Ala Val Ile Ser Tyr Asp Gly Thr Asn Lys Tyr Phe Ala Asp Ser Val
426 50 55 60
430 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
432 65 70 75 80
436 Leu Gln Met Thr Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Phe Cys

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438      85      90      95
442 Ala Asn Leu Arg Gly Glu Val Thr Arg Arg Ala Ser Val Pro Leu Asp
444      100      105      110
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480 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe Ser Ser Tyr
482 20 25 30
486 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
488 35 40 45
492 Ser Val Ile Ser Tyr Asp Gly His His Lys Asn Tyr Ala Asp Ser Val
494 50 55 60
498 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Lys Thr Leu Tyr
500 65 70 75 80
504 Leu Gln Met Asn Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys
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548 Ser Leu Arg Leu Ser Cys Val Val Ser Gly Phe Thr Phe Asn Asn Tyr
550 20 25 30
554 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
556 35 40 45
560 Ala Val Ile Trp Phe Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val
562 50 55 60
566 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
568 65 70 75 80
572 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
574 85 90 95
578 Ala Arg Glu Asn Gln Ile Lys Leu Trp Ser Arg Tyr Leu Tyr Tyr Phe
580 100 105 110

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618      20           25           30
622 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
624      35           40           45
628 Ala Val Ile Trp Phe Asp Gly Ser Asn Lys Glu Tyr Ala Asp Ser Val
630      50           55           60
634 Lys Gly Arg Phe Thr Val Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
636   65           70           75           80
640 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
642      85           90           95
646 Ala Arg Glu Glu Val Val Arg Gly Val Ile Leu Trp Ser Arg Lys Phe
648      100          105          110
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680   1           5           10           15
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686      20           25           30
690 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
692      35           40           45
696 Ala Asp Ile Trp Phe Asp Gly Ser Asn Lys Asp Tyr Ala Asp Ser Val
698      50           55           60
702 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
704   65           70           75           80
708 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
710      85           90           95
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VERIFICATION SUMMARY

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